

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/580,906
Source: 1fwsp
Date Processed by STIC: 6/8/06

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

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- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

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Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
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Revised 01/10/06



IFWP

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/580,906

DATE: 06/08/2006
TIME: 10:08:05

Input Set : A:\Sequence Listing for RECEPTOR FUNCTION REGULATING

AGENT.txt

Output Set: N:\CRF4\06082006\J580906.raw

3 <110> APPLICANT: FUKATSU et al.
5 <120> TITLE OF INVENTION: RECEPTOR FUNCTION REGULATING AGENT
7 <130> FILE REFERENCE: 20039.0005USWO
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/580,906
10 <141> CURRENT FILING DATE: 2006-05-26
12 <150> PRIOR APPLICATION NUMBER: PCT/JP2004/017996
13 <151> PRIOR FILING DATE: 2004-11-26
15 <150> PRIOR APPLICATION NUMBER: JP 2003-394848
16 <151> PRIOR FILING DATE: 2003-11-26
18 <160> NUMBER OF SEQ ID NOS: 20
20 <170> SOFTWARE: PatentIn Version 3.1
22 <210> SEQ ID NO: 1
23 <211> LENGTH: 361
24 <212> TYPE: PRT
25 <213> ORGANISM: Human

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W--> 26 <400> SEQUENCE: 1
28 Met Ser Pro Glu Cys Ala Arg Ala Ala Gly Asp Ala Pro Leu Arg Ser
29 . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
29 5 10 15
30 Leu Glu Gln Ala Asn Arg Thr Arg Phe Pro Phe Phe Ser Asp Val Lys
31 . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
31 20 25 30
32 Gly Asp His Arg Leu Val Leu Ala Ala Val Glu Thr Thr Val Leu Val
33 . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
33 35 40 45
34 Leu Ile Phe Ala Val Ser Leu Leu Gly Asn Val Cys Ala Leu Val Leu
35 . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
35 50 55 60
36 Val Ala Arg Arg Arg Arg Gly Ala Thr Ala Cys Leu Val Leu Asn
37 . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
37 65 70 75 80
38 Leu Phe Cys Ala Asp Leu Leu Phe Ile Ser Ala Ile Pro Leu Val Leu
39 . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
39 85 90 95
40 Ala Val Arg Trp Thr Glu Ala Trp Leu Leu Gly Pro Val Ala Cys His
41 . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
41 100 105 110
42 Leu Leu Phe Tyr Val Met Thr Leu Ser Gly Ser Val Thr Ile Leu Thr
43 . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
43 115 120 125
44 Leu Ala Ala Val Ser Leu Glu Arg Met Val Cys Ile Val His Leu Gln
45 . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
45 130 135 140
46 Arg Gly Val Arg Gly Pro Gly Arg Arg Ala Arg Ala Val Leu Leu Ala
47 . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
47 145 150 155 160
48 Leu Ile Trp Gly Tyr Ser Ala Val Ala Ala Leu Pro Leu Cys Val Phe
49 . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
49 165 170 175
50 Phe Arg Val Val Pro Gln Arg Leu Pro Gly Ala Asp Gln Glu Ile Ser
51 . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
51 . 180 185 190
52 Ile Cys Thr Leu Ile Trp Pro Thr Ile Pro Gly Glu Ile Ser Trp Asp
53 . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
53 195 200 205
54 Val Ser Phe Val Thr Leu Asn Phe Leu Val Pro Gly Leu Val Ile Val

```

pp 4, 6
Does Not Comply
Corrected Diskette Needed

RAW SEQUENCE LISTING
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Input Set : A:\Sequence Listing for RECEPTOR FUNCTION REGULATING
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Output Set: N:\CRF4\06082006\J580906.raw

55	210	215	220
56	Ile Ser Tyr Ser Lys Ile Leu Gln Ile Thr Lys Ala Ser Arg Lys Arg		
57	225	230	235
58	Leu Thr Val Ser Leu Ala Tyr Ser Glu Ser His Gln Ile Arg Val Ser		240
59	245	250	255
60	Gln Gln Asp Phe Arg Leu Phe Arg Thr Leu Phe Leu Leu Met Val Ser		
61	260	265	270
62	Phe Phe Ile Met Trp Ser Pro Ile Ile Thr Ile Leu Leu Ile Leu		
63	275	280	285
64	Ile Gln Asn Phe Lys Gln Asp Leu Val Ile Trp Pro Ser Leu Phe Phe		
65	290	295	300
66	Trp Val Val Ala Phe Thr Phe Ala Asn Ser Ala Leu Asn Pro Ile Leu		
67	305	310	315
68	Tyr Asn Met Thr Leu Cys Arg Asn Glu Trp Lys Lys Ile Phe Cys Cys		320
69	325	330	335
70	Phe Trp Phe Pro Glu Lys Gly Ala Ile Leu Thr Asp Thr Ser Val Lys		
71	340	345	350
72	Arg Asn Asp Leu Ser Ile Ile Ser Gly		
73	355	360	
74	<210> SEQ ID NO: 2		
75	<211> LENGTH: 1083		
76	<212> TYPE: DNA		
77	<213> ORGANISM: Human		
W-->	78 <400> SEQUENCE: 2		
79	atgccccctg aatgcgcgcg ggcagcgggc gacgcgcct tgcgcaagcc ggagcaagcc	60	
80	aaccgcaccc gctttccctt cttctccgac gtcaaggcg accaccggct ggtgtggcc	120	
81	gcgggtggaga caaccgtgct ggtgctcatc tttgcagtgt cgctgctggg caacgtgtgc	180	
82	gccctgggtgc tgggtggcg cgacgcacgc cgccggcgca ctgcctgcct ggtactcaac	240	
83	ctcttctgct cggacctgct cttcatcagc gctatccctc tgggtgctggc cgtgcgctgg	300	
84	actgaggccct ggctgctggg ccccgttgcc tgccacctgc tcttctacgt gatgaccctg	360	
85	agccgcagcgc tcaccatcct cacgctggcc gcggtcagcc tggagcgcata ggtgtgcata	420	
86	gtgcacctgc agcgcgcgt ggggggtctt gggcgccgg cgccggcagt gctgctggcg	480	
87	ctcatctggg gctattcggc ggtgcccgcct ctgcctctct gcgtcttctt ccgagtcgtc	540	
88	ccgcaacggc tccccggcgc cgaccaggaa atttcgattt gcacactgat ttggcccacc	600	
89	attcctggag agatctcgta ggtatgtctt tttgttactt tgaacttctt ggtgccagga	660	
90	ctggtcattt tgatcagttt ctccaaaattt ttacagatca caaaggcatc aaggaagagg	720	
91	ctcacggtaa gcctggcta ctcggagagc caccagatcc gcgtgtccca gcaggacttc	780	
92	cggctttcc gcaccctttt cttccatcgat gtctcattt tcatcatgtg gagccccatc	840	
93	atcatcacca tccttcat cctgatccag aacctcaagc aagacctggt catctggccg	900	
94	tccctttct tctgggtggt ggcttcaca tttgctaattt cagccctaaa ccccatcctc	960	
95	tacaacatga cactgtgcag gaatgagtgg aaaaaattt tttgctgctt ctggcccacca	1020	
96	gaaaaggag ccatttaac agacacatct gtcaaaaagaa atgacttgta gattatttct	1080	
97	ggc	1083	
98	<210> SEQ ID NO: 3		
99	<211> LENGTH: 361		
100	<212> TYPE: PRT		
101	<213> ORGANISM: Mouse		
W-->	102 <400> SEQUENCE: 3		
103	Met Ser Pro Glu Cys Ala Gln Thr Thr Gly Pro Gly Pro Ser His Thr		

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Output Set: N:\CRF4\06082006\J580906.raw

```

104      5          10          15
105 Leu Asp Gln Val Asn Arg Thr His Phe Pro Phe Ser Asp Val Lys
106      20         25          30
107 Gly Asp His Arg Leu Val Leu Ser Val Val Glu Thr Thr Val Leu Gly
108      35         40          45
109 Leu Ile Phe Val Val Ser Leu Leu Gly Asn Val Cys Ala Leu Val Leu
110      50         55          60
111 Val Ala Arg Arg Arg Arg Gly Ala Thr Ala Ser Leu Val Leu Asn
112      65         70          75          80
113 Leu Phe Cys Ala Asp Leu Leu Phe Thr Ser Ala Ile Pro Leu Val Leu
114      85         90          95
115 Val Val Arg Trp Thr Glu Ala Trp Leu Leu Gly Pro Val Val Cys His
116      100        105         110
117 Leu Leu Phe Tyr Val Met Thr Met Ser Gly Ser Val Thr Ile Leu Thr
118      115        120         125
119 Leu Ala Ala Val Ser Leu Glu Arg Met Val Cys Ile Val Arg Leu Arg
120      130        135         140
121 Arg Gly Leu Ser Gly Pro Gly Arg Arg Thr Gln Ala Ala Leu Leu Ala
122 145      150        155         160
123 Phe Ile Trp Gly Tyr Ser Ala Leu Ala Ala Leu Pro Leu Cys Ile Leu
124      165        170         175
125 Phe Arg Val Val Pro Gln Arg Leu Pro Gly Gly Asp Gln Glu Ile Pro
126      180        185         190
127 Ile Cys Thr Leu Asp Trp Pro Asn Arg Ile Gly Glu Ile Ser Trp Asp
128      195        200         205
129 Val Phe Phe Val Thr Leu Asn Phe Leu Val Pro Gly Leu Val Ile Val
130      210        215         220
131 Ile Ser Tyr Ser Lys Ile Leu Gln Ile Thr Lys Ala Ser Arg Lys Arg
132 225      230        235         240
133 Leu Thr Leu Ser Leu Ala Tyr Ser Glu Ser His Gln Ile Arg Val Ser
134      245        250         255
135 Gln Gln Asp Tyr Arg Leu Phe Arg Thr Leu Phe Leu Leu Met Val Ser
136      260        265         270
137 Phe Phe Ile Met Trp Ser Pro Ile Ile Ile Thr Ile Leu Leu Ile Leu
138      275        280         285
139 Ile Gln Asn Phe Arg Gln Asp Leu Val Ile Trp Pro Ser Leu Phe Phe
140      290        295         300
141 Trp Val Val Ala Phe Thr Phe Ala Asn Ser Ala Leu Asn Pro Ile Leu
142 305      310        315         320
143 Tyr Asn Met Ser Leu Phe Arg Asn Glu Trp Arg Lys Ile Phe Cys Cys
144      325        330         335
145 Phe Phe Phe Pro Glu Lys Gly Ala Ile Phe Thr Asp Thr Ser Val Arg
146      340        345         350
147 Arg Asn Asp Leu Ser Val Ile Ser Ser
148      355        360
149 <210> SEQ ID NO: 4
150 <211> LENGTH: 1083
151 <212> TYPE: DNA
152 <213> ORGANISM: Mouse

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RAW SEQUENCE LISTING
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Input Set : A:\Sequence Listing for RECEPTOR FUNCTION REGULATING
AGENT.txt
Output Set: N:\CRF4\06082006\J580906.raw

W--> 153 <400> SEQUENCE: 4

154	atgtccctg	agtgtcaca	gacgacgggc	cctggcccct	cgcacaccct	ggaccaagtc	60
155	aatcgaccc	acttccctt	cttctcgat	gtcaaggcg	accaccgggtt	ggtgttgagc	120
156	gtcgtggaga	ccaccgttct	ggggctcatc	tttgcgtct	cactgctggg	caacgtgtgt	180
157	gctcttagtgc	tggtggcgcg	ccgtcgccgc	cgtggggcga	cagccagcc	ggtgctcaac	240
158	ctcttctcg	cggatttgc	tttaccaggc	gccatccctc	tagtgcctgt	cgtgcgtgg	300
159	actgaggcct	ggctgttggg	gcccgtcgtc	tgcacacctgc	tcttctacgt	gatgacaatg	360
160	agcggcagcg	tcacgatcct	cacactggcc	gcgggtcagcc	tggagcgcac	ggtgtgcac	420
161	gtgcgcctcc	ggcgcggcctt	gagcggcccg	gggcggcgg	ctcaggcggc	actgctggct	480
162	ttcatatggg	gttactcggc	gtcgcgcgc	ctgcgcctct	gcatcttgc	ccgcgtggc	540
163	ccgcagcgcc	ttccccggcg	ggaccaggaa	attccgattt	gcacattgga	ttggcccaac	600
164	cgcataaggag	aaatctcatg	gatgtgttt	tttgcgtactt	tgaacttcct	ggtgcgggga	660
165	ctggtcattt	tgatcagttt	ctccaaaattt	ttacagatca	cgaaagcatc	gcggaagagg	720
166	cttacgctga	gcttggcata	ctctgagagc	caccagatcc	gagtgtccca	acaagactac	780
167	cgaactcttcc	gcacgcttct	cctgctcatg	gttcgcctct	tcatcatgtg	gagtcacatc	840
168	atcatcacca	tcctcctcat	tttgatccaa	aactccggc	aggacactgtt	catctggcca	900
169	tccctttct	tctgggttgt	ggccttcacg	tttgcctact	ctgcctaaa	ccccataactg	960
170	tacaacatgt	cgctgttcag	gaacgaatgg	aggaagattt	tttgcgtctt	ctttttcca	1020
171	gagaagggag	ccattttac	agacacgtct	gtcaggcga	atgacttgc	tgttatttcc	1080
172	agc						1083

173 <210> SEQ ID NO: 5

174 <211> LENGTH: 20

175 <212> TYPE: DNA

176 <213> ORGANISM: Artificial Sequence

W--> 177 <220> FEATURE:

W--> 178 <223> OTHER INFORMATION:

W--> 178 <400> SEQUENCE: 5

179 gctgtggcat gcttttaaac

20

180 <210> SEQ ID NO: 6

181 <211> LENGTH: 20

182 <212> TYPE: DNA

183 <213> ORGANISM: Artificial Sequence

W--> 184 <220> FEATURE:

W--> 185 <223> OTHER INFORMATION:

W--> 185 <400> SEQUENCE: 6

186 cgctgtggat gtctatttgc

20

187 <210> SEQ ID NO: 7

188 <211> LENGTH: 30

189 <212> TYPE: DNA

190 <213> ORGANISM: Artificial Sequence

W--> 191 <220> FEATURE:

W--> 192 <223> OTHER INFORMATION:

W--> 192 <400> SEQUENCE: 7

193 agttcatttc cagtaccctc catcagtggc

30

194 <210> SEQ ID NO: 8

195 <211> LENGTH: 361

196 <212> TYPE: PRT

197 <213> ORGANISM: Rat

W--> 198 <400> SEQUENCE: 8

) see p. 6 for error explanation'

This error appears in other sequences, too

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**Input Set : A:\Sequence Listing for RECEPTOR FUNCTION REGULATING
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```

199 Met Ser Pro Glu Cys Ala Gln Thr Thr Gly Pro Gly Pro Ser Arg Thr
200           5          10          15
201 Pro Asp Gln Val Asn Arg Thr His Phe Pro Phe Phe Ser Asp Val Lys
202           20         25          30
203 Gly Asp His Arg Leu Val Leu Ser Val Leu Glu Thr Thr Val Leu Gly
204           35         40          45
205 Leu Ile Phe Val Val Ser Leu Leu Gly Asn Val Cys Ala Leu Val Leu
206           50         55          60
207 Val Val Arg Arg Arg Arg Gly Ala Thr Val Ser Leu Val Leu Asn
208           65         70          75          80
209 Leu Phe Cys Ala Asp Leu Leu Phe Thr Ser Ala Ile Pro Leu Val Leu
210           85         90          95
211 Val Val Arg Trp Thr Glu Ala Trp Leu Leu Gly Pro Val Val Cys His
212           100        105         110
213 Leu Leu Phe Tyr Val Met Thr Met Ser Gly Ser Val Thr Ile Leu Thr
214           115        120         125
215 Leu Ala Ala Val Ser Leu Glu Arg Met Val Cys Ile Val Arg Leu Arg
216           130        135         140
217 Arg Gly Leu Ser Gly Pro Gly Arg Arg Thr Gln Ala Ala Leu Leu Ala
218           145        150         155         160
219 Phe Ile Trp Gly Tyr Ser Ala Leu Ala Ala Leu Pro Leu Cys Ile Leu
220           165        170         175
221 Phe Arg Val Val Pro Gln Arg Leu Pro Gly Gly Asp Gln Glu Ile Pro
222           180        185         190
223 Ile Cys Thr Leu Asp Trp Pro Asn Arg Ile Gly Glu Ile Ser Trp Asp
224           195        200         205
225 Val Phe Phe Val Thr Leu Asn Phe Leu Val Pro Gly Leu Val Ile Val
226           210        215         220
227 Ile Ser Tyr Ser Lys Ile Leu Gln Ile Thr Lys Ala Ser Arg Lys Arg
228           225        230         235         240
229 Leu Thr Leu Ser Leu Ala Tyr Ser Glu Ser His Gln Ile Arg Val Ser
230           245        250         255
231 Gln Gln Asp Tyr Arg Leu Phe Arg Thr Leu Phe Leu Leu Met Val Ser
232           260        265         270
233 Phe Phe Ile Met Trp Ser Pro Ile Ile Thr Ile Leu Leu Ile Leu
234           275        280         285
235 Ile Gln Asn Phe Arg Gln Asp Leu Val Ile Trp Pro Ser Leu Phe Phe
236           290        295         300
237 Trp Val Val Ala Phe Thr Phe Ala Asn Ser Ala Leu Asn Pro Ile Leu
238           305        310         315         320
239 Tyr Asn Met Ser Leu Phe Arg Ser Glu Trp Arg Lys Ile Phe Cys Cys
240           325        330         335
241 Phe Phe Phe Pro Glu Lys Gly Ala Ile Phe Thr Glu Thr Ser Ile Arg
242           340        345         350
243 Arg Asn Asp Leu Ser Val Ile Ser Thr
244           355        360
245 <210> SEQ ID NO: 9
246 <211> LENGTH: 1083
247 <212> TYPE: DNA

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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 06/08/2006
PATENT APPLICATION: US/10/580,906 TIME: 10:08:06

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Use of <220> Feature(NEW RULES):

error explanation

Sequence(s) are missing the <220> Feature and associated headings.

Use of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial Sequence" or "Unknown". Please explain source of genetic material in <220> to <223> section (See "Federal Register," 6/01/98, Vol. 63, No. 104, pp.29631-32)
(Sec.1.823 of new Rules)

Seq#:5,6,7,10,11,12,13,14,15,16,17,18,19,20

VERIFICATION SUMMARY **DATE: 06/08/2006**
PATENT APPLICATION: US/10/580,906 **TIME: 10:08:06**

Input Set : A:\Sequence Listing for RECEPTOR FUNCTION REGULATING
AGENT.txt
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L:9 M:270 C: Current Application Number differs, Replaced Current Application Number
L:26 M:283 W: Missing Blank Line separator, <400> field identifier
L:78 M:283 W: Missing Blank Line separator, <400> field identifier
L:102 M:283 W: Missing Blank Line separator, <400> field identifier
L:153 M:283 W: Missing Blank Line separator, <400> field identifier
L:177 M:283 W: Missing Blank Line separator, <220> field identifier
L:178 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:5, <213>
ORGANISM:Artificial Sequence
L:178 M:283 W: Missing Blank Line separator, <400> field identifier
L:178 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:5,Line#:178
L:184 M:283 W: Missing Blank Line separator, <220> field identifier
L:185 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:6, <213>
ORGANISM:Artificial Sequence
L:185 M:283 W: Missing Blank Line separator, <400> field identifier
L:185 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:6,Line#:185
L:191 M:283 W: Missing Blank Line separator, <220> field identifier
L:192 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:7, <213>
ORGANISM:Artificial Sequence
L:192 M:283 W: Missing Blank Line separator, <400> field identifier
L:192 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:7,Line#:192
L:198 M:283 W: Missing Blank Line separator, <400> field identifier
L:249 M:283 W: Missing Blank Line separator, <400> field identifier
L:273 M:283 W: Missing Blank Line separator, <220> field identifier
L:274 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:10, <213>
ORGANISM:Artificial Sequence
L:274 M:283 W: Missing Blank Line separator, <400> field identifier
L:274 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:10,Line#:274
L:280 M:283 W: Missing Blank Line separator, <220> field identifier
L:281 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:11, <213>
ORGANISM:Artificial Sequence
L:281 M:283 W: Missing Blank Line separator, <400> field identifier
L:281 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:11,Line#:281
L:287 M:283 W: Missing Blank Line separator, <220> field identifier
L:288 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:12, <213>
ORGANISM:Artificial Sequence
L:288 M:283 W: Missing Blank Line separator, <400> field identifier
L:288 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:12,Line#:288
L:294 M:283 W: Missing Blank Line separator, <220> field identifier
L:295 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:13, <213>
ORGANISM:Artificial Sequence
L:295 M:283 W: Missing Blank Line separator, <400> field identifier
L:295 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:13,Line#:295
L:301 M:283 W: Missing Blank Line separator, <220> field identifier
L:302 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:14, <213>
ORGANISM:Artificial Sequence
L:302 M:283 W: Missing Blank Line separator, <400> field identifier
L:302 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:14,Line#:302
L:308 M:283 W: Missing Blank Line separator, <220> field identifier
L:309 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:15, <213>
ORGANISM:Artificial Sequence
L:309 M:283 W: Missing Blank Line separator, <400> field identifier
L:309 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:15,Line#:309

L:315 M:283 W: Missing Blank Line separator, <220> field identifier
L:316 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:16, <213>
ORGANISM:Artificial Sequence
L:316 M:283 W: Missing Blank Line separator, <400> field identifier
L:316 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:16,Line#:316
L:322 M:283 W: Missing Blank Line separator, <220> field identifier

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/580,906

DATE: 06/08/2006

TIME: 10:08:06

Input Set : A:\Sequence Listing for RECEPTOR FUNCTION REGULATING

AGENT.txt

Output Set: N:\CRF4\06082006\J580906.raw

L:323 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:17, <213>
ORGANISM:Artificial Sequence

L:323 M:283 W: Missing Blank Line separator, <400> field identifier

L:323 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:17,Line#:323

L:329 M:283 W: Missing Blank Line separator, <220> field identifier

L:330 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:18, <213>

ORGANISM:Artificial Sequence

L:330 M:283 W: Missing Blank Line separator, <400> field identifier

L:330 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:18,Line#:330

L:336 M:283 W: Missing Blank Line separator, <220> field identifier

L:337 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:19, <213>

ORGANISM:Artificial Sequence

L:337 M:283 W: Missing Blank Line separator, <400> field identifier

L:337 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:19,Line#:337

L:343 M:283 W: Missing Blank Line separator, <220> field identifier

L:344 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:20, <213>

ORGANISM:Artificial Sequence

L:344 M:283 W: Missing Blank Line separator, <400> field identifier

L:344 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:20,Line#:344